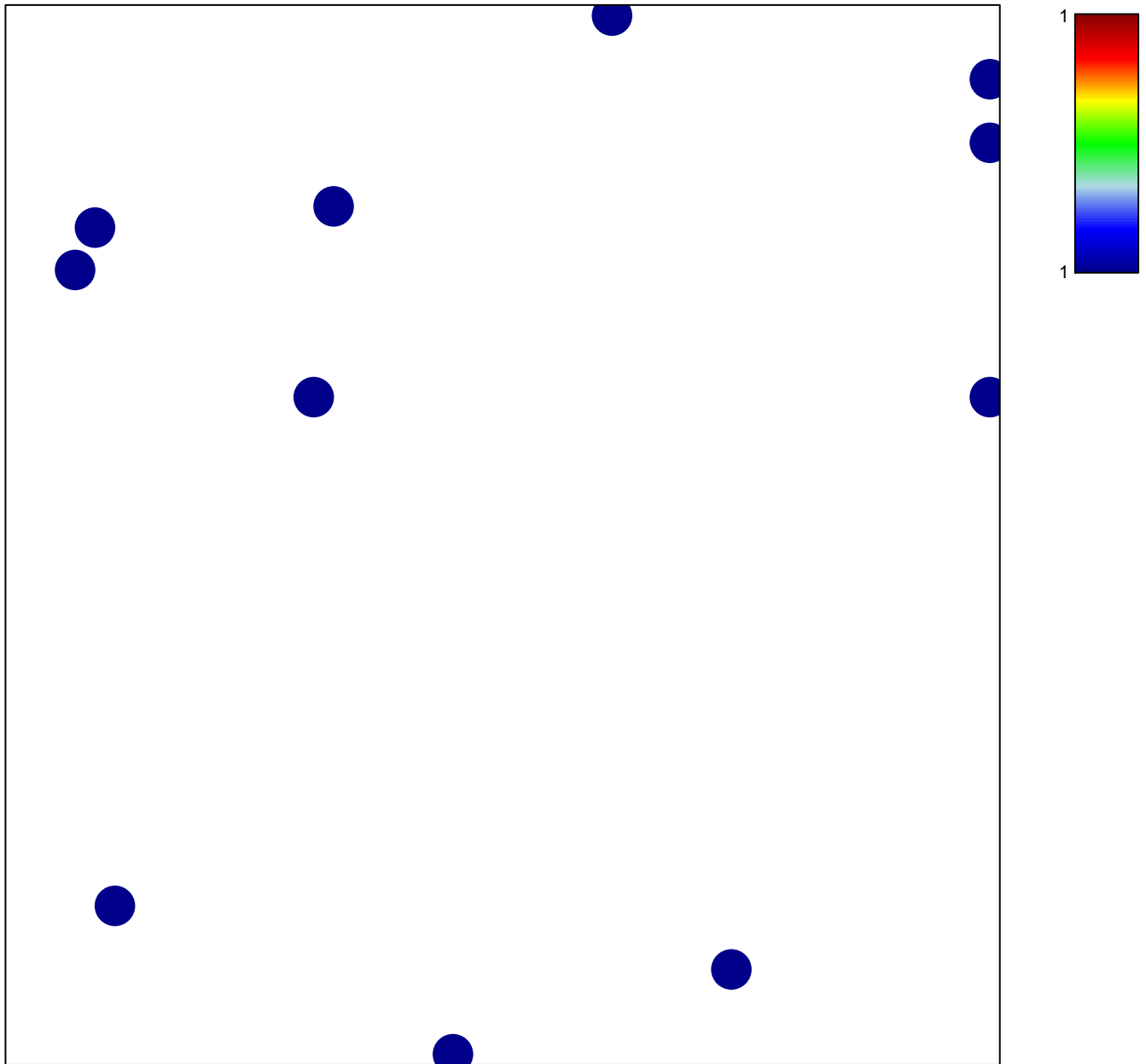
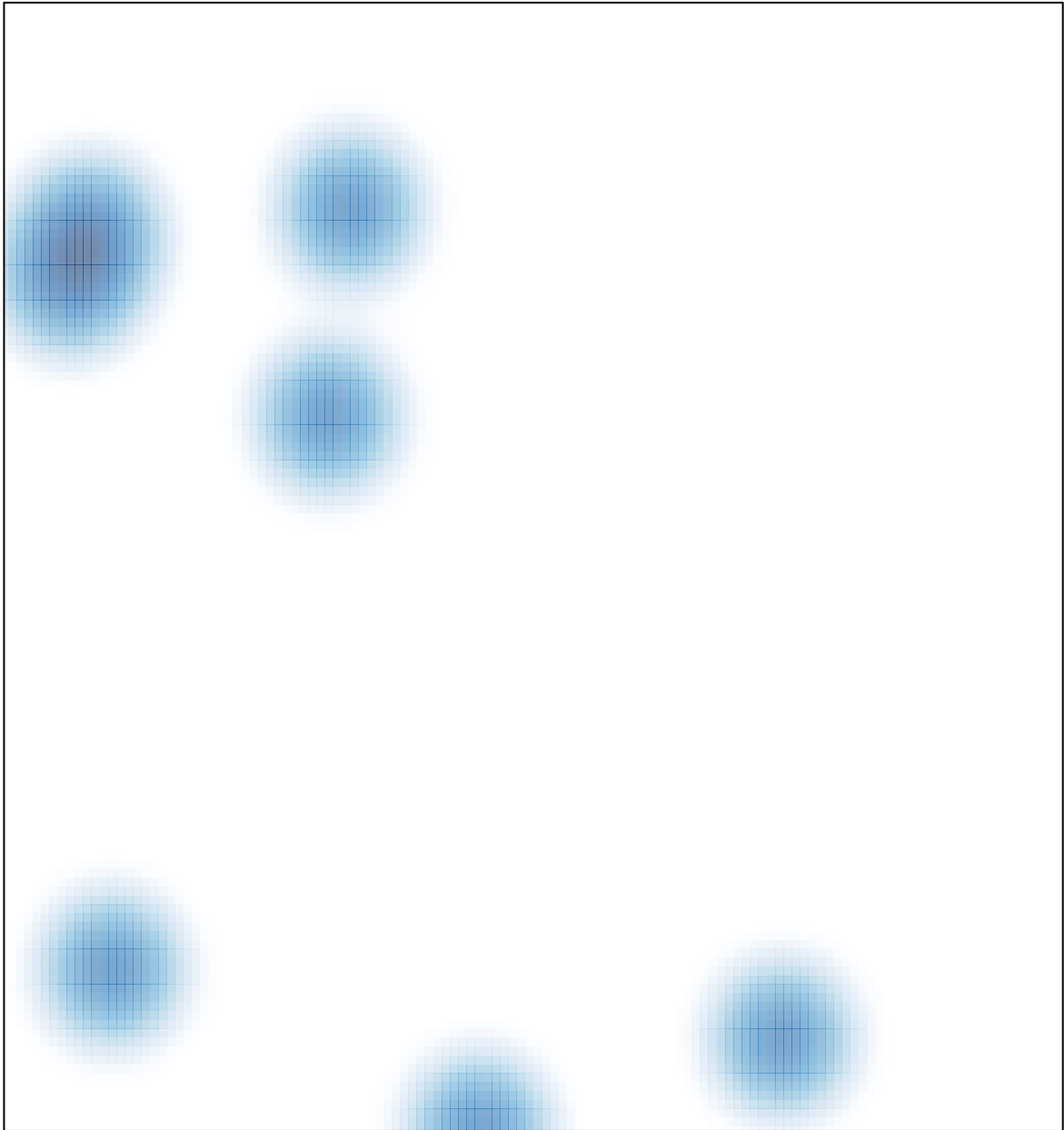


chaperone-mediated protein complex assembly



features = 11
chi-square p = 0.83

chaperone-mediated protein complex assembly



features = 11 , max = 1